

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING** **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/562,627  
Source: IFWP  
Date Processed by STIC: 1/10/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

## Raw Sequence Listing Error Summary

### ERROR DETECTED

### SUGGESTED CORRECTION

SERIAL NUMBER: 10/562,627

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1      Wrapped Nucleics  
    Wrapped Aminos     The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2      Invalid Line Length     The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
- 3      Misaligned Amino  
    Numbering     The numbering under each 5<sup>th</sup> amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
- 4      Non-ASCII     The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. Please **ensure your subsequent submission is saved in ASCII text**.
- 5      Variable Length     Sequence(s)      contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6      PatentIn 2.0  
    "bug"     A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)     . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
- 7      Skipped Sequences  
    (OLD RULES)     Sequence(s)      missing. If intentional, please insert the following lines for **each** skipped sequence:  
                          (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
                          (i)     SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
                          (xi)  SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
                          This sequence is intentionally skipped  
  
                          Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
- 8      Skipped Sequences  
    (NEW RULES)     Sequence(s)      missing. If **intentional**, please insert the following lines for **each** skipped sequence.  
                          <210> sequence id number  
                          <400> sequence id number  
                          000
- 9      Use of n's or Xaa's  
    (NEW RULES)     Use of n's and/or Xaa's have been detected in the Sequence Listing.  
                          Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.  
                          In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
- 10     Invalid <213>  
    Response     Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
- 11     Use of <220>     Sequence(s)      missing the <220> "Feature" and associated numeric identifiers and responses.  
                          Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
                          (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12     PatentIn 2.0  
    "bug"     Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13     Misuse of n/Xaa     "**n**" can **only** represent a single nucleotide; "**Xaa**" can **only** represent a single amino acid



IFWP

## RAW SEQUENCE LISTING

DATE: 01/10/2006

PATENT APPLICATION: US/10/562,627

TIME: 09:00:14

Input Set : N:\DA\PTO.DA.txt

Output Set: N:\CRF4\01102006\J562627.raw

*pp 1-5*  
**Does Not Comply  
 Corrected Diskette Needed**

*see item 2 on Enr summary sheet*

3 <110> APPLICANT: CHOE, Mu-Hyeon  
 4 CHOI, Seong-Hyeok  
 5 LEE, Yong-Chan  
 6 KWON, Hye-Won  
 7 WON, Jae-Seon  
 8 YU, Mi-Hyun  
 9 SONG, Jeong-Hwa  
 10 KIM, Yong-Jae

12 <120> **TITLE OF INVENTION:** The Dimer of Chimeric Recombinant Binding Domain-Functional  
 Group

13 Fusion formed via Disulfide-bond-bridge and The Process For Producing The Same

15 <130> FILE REFERENCE: 428.1060

C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/562,627

C--> 17 <141> CURRENT FILING DATE: 2005-12-22

17 <150> PRIOR APPLICATION NUMBER: PCT/KR2004/001595

18 <151> PRIOR FILING DATE: 2004-06-30

20 <150> PRIOR APPLICATION NUMBER: KR2003-0043599

21 <151> PRIOR FILING DATE: 2003-06-30

23 <160> NUMBER OF SEQ ID NOS: 12

25 <170> SOFTWARE: KopatentIn 1.71

27 <210> SEQ ID NO: 1

28 <211> LENGTH: 1749

29 <212> TYPE: DNA

30 <213> ORGANISM: pmc74 plasmid coding sequence

32 <400> SEQUENCE: 1

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35	ctctcctgtg caacctctgg attcactttc agtgactatt acatgtattg ggttcgccag	120
37	actccagaga agaggctgga gtgggtcgca tacattagta atgatgatag ttccgcccgt	180
39	tattcagaca ctgtaaaggg ccggttcacc atctccagag acaatgccag gaacaccctc	240
41	tacctgcaaa tgagccgtct gaagtctgag gacacagcca tatattcctg tgcaagagga	300
43	ctggcctggg gagcctggtt tgcttactgg ggccaaggga ctctggtcac tgtctctgca	360
45	gccaaaacga ccccccatc tgtctatcca ctggcccctg gatctgctgc ccaaactaac	420
47	tccatggtga ccctgggatg cctggtcaag ggctatttcc ctgagccagt gacagtgacc	480
49	tggaaactctg gatccctgtc cagcgggtgtg cacaccttcc cagctgtcct gcagtctgac	540
51	ctctacactc tgagcagctc agtgactgtc ccctccagca cctggcccag cgagaccgtc	600
53	acctgcaacg ttgcccaccc ggccagcagc accaagggtg acaagaaaat tgtgcccagg	660
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57	ctggccgcgc tgaccgcgca ccaggcttgc cacctgccgc tggagacttt caccggtcat	780
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61	gccctctacc tggcggcgcg gctgtcgtgg aaccaggtcg accaggtgat ccgcaacgcc	900
63	ctggccagcc ccggcagcgg cggcgacctg ggcgaaagca tccgcgagca gccggagcag	960
65	gcccgctctg ccctgaccct ggccgcccgc gagagcgagc gcttcgtccg gcagggcacc	1020
67	ggcaacgacg aggccggcgc ggccaacggc ccggcgagca gcggcgacgc cctgctggag	1080
69	cgcaactatc ccactggcgc ggagttcctc ggcgacggcg gcgacgtcag cttcagcacc	1140

*Invalid <213> response see item 10 on Enr summary sheet.*

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Input Set : N:\DA\PTO.DA.txt

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71 cgcggcacgc agaactggac ggtggagcgg ctgctccagg cgcaccgcca actggaggag 1200
73 cgcggcctatg tgttcgtcgg ctaccacggc accttcctcg aagcggcgca aagcatcgtc 1260
75 ttcggcgggg tgcgcgcgcg cagccaggac ctgcagcgca tctggcgcgg tttctatatc 1320
77 gccggcgatc cggcgctggc ctacggctac gcccaggacc aggaaccgga cgcacgcggc 1380
79 cggatccgca acggtgccct gctgcgggtc tatgtgccgc gctcgagcct gccgggcttc 1440
81 taccgcacca gcctgaccct ggccgcgcgc gagggcgcgg gcgaggtcga acggctgatc 1500
83 ggccatccgc tgccgctgcg cctggacgcc atcaccggcc ccgaggagga aggcgggcgc 1560
85 ctggagacca ttctcggtg gccgctggcc gagcgaccgc tggtgattcc ctccggcgatc 1620
87 cccaccgacc cgcgcaacgt cggcggcgac ctgcaccgct ccagcatccc cgacaaggaa 1680
89 caggcgatca gcgccctgcc ggactacgcc agccagcccg gcaaacgcc gcgcgaggac 1740
91 ctgaagtaa 1749
94 <210> SEQ ID NO: 2
95 <211> LENGTH: 1764
96 <212> TYPE: DNA
97 <213> ORGANISM: PMH21 plasmid coding sequence
98 <400> SEQUENCE: 2
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102 ctctcctgtg caacctctgg attcactttc agtgactatt acatgtattg ggttcgccag 120
104 actccagaga agaggctgga gtgggtcgca tacattagta atgatgatag ttccgccgct 180
106 tattcagaca ctgtaaaggc ccggttcacc atctccagag acaatgccag gaacaccctc 240
108 tacctgcaaa tgagccgtct gaagtctgag gacacagcca tatattcctg tgcaagagga 300
110 ctggcctggg gagcctgggt tgcttactgg ggccaaggga ctctgggtcac tgtctctgca 360
112 gccaaaacga ccccccatc tgtctatcca ctggcccctg gatctgctgc ccaaactaac 420
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120 acctgcaacg ttgcccaccc ggccagcagc accaagggtg acaagaaaat tgtgcccagg 660
122 gattgtggta gtaagccttg cataagtaca aaagcttctg gtggtggcgg atctggagggt 720
124 cccgagggcg gcagcctggc cgcgctgacc gcgcaccagg cttgccacct gccgctggag 780
126 actttcaccc gtcctgcgca gccgcgcggc tgggaacaac tggagcagtg cggctatccg 840
128 gtgcagcggc tggtcgccct ctacctggcg gcgcggctgt cgtggaacca ggtcgaccag 900
130 gtgatccgca acgcccctggc cagccccggc agcggcgggc acctgggcca agcgatccgc 960
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136 gacgccctgc tggagcgcaa ctatccact ggcgcgaggt tcctcggcga cggcggcgac 1140
138 gtcagcttca gcaccgcgg cagcgagaac tggacgggtg agcggctgct ccaggcgcac 1200
140 cgccaaactg aggagcgcg ctatgtgttc gtcggctacc acggcacctt cctcgaagcg 1260
142 gcgcaaagca tcgtcttcgg cggggtgcgc gcgcgcagcc aggaacctga cgcgatctgg 1320
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146 cccgacgcac gcggccggat ccgcaacggg gccctgctgc ggggtctatgt gccgcgctcg 1440
148 agcctgccgg gcttctaccg caccagcctg accctggccg cgccggaggc ggcgggcgag 1500
150 gtcgaacggc tgatcggccca tccgctgccg ctgcgcctgg acgccatcac cggccccgag 1560
152 gaggaaggcg ggcgcctgga gaccattctc ggctggccgc tggccgagcg caccgtgggtg 1620
154 attccctcgg cgatccccac cgaccgcgc aacgtcggcg gcgacctga cccgtccagc 1680
156 atccccgaca aggaacaggc gatcagcgcc ctgcccggact acgccagcca gcccggcaaa 1740
158 ccgcccgcgc aggaacctgaa gtaa
161 <210> SEQ ID NO: 3
162 <211> LENGTH: 1749
163 <212> TYPE: DNA

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## RAW SEQUENCE LISTING

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Output Set: N:\CRF4\01102006\J562627.raw

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164 <213> ORGANISM: pCE2 plasmid coding sequence
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169 ctctcctgtg caacctctgg attcactttc agtgactatt acatgtattg ggttcgccag      120
171 actccagaga agaggctgga gtgggtcgca tacattagta atgatgatag ttccgcccgt      180
173 tattcagaca ctgtaaaggg ccggttcacc atctccagag acaatgccag gaacaccctc      240
175 tacctgcaaa tgagccgtct gaagtctgag gacacagcca tatattcctg tgcaagagga      300
177 ctggcctggg gagcctgggt tgcttactgg ggccaaggga ctctggtcac tgtctctgca      360
179 gccaaaacga cacccccata tgtctatcca ctggcccctg gatctgctgc ccaaactaac      420
181 tccatggtga ccctgggatg cctgggcaag ggctatttcc ctgagccagt gacagtgacc      480
183 tggaactctg gatccctgtc cagcgggtgt cacaccttcc cagctgtcct gcagtctgac      540
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187 acctgcaacg ttgcccaccc ggccagcagc accaagggtg acaagaaaat tgtgcccagg      660
189 gatttgtgga gtaagccttg cataagtaca aaagcttccg gagggtcccga gggcggcagc      720
191 ctggccgcgc tgaccgcgca ccaggcttgc cacctgccgc tggagacttt caccgcgtcat      780
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199 gccgctctgg ccctgaccct ggccgcgccg gagagcgagc gcttcgtccg gcagggcacc      1020
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207 cgcggtctat gtgtcgctcg ctaccacggc accttctctg aagcggcgca aagcatcgct      1260
209 ttcggcgggg tgcgcgcgcg cagccaggac ctcgacgcga tctggcgcgg ttctatatatc      1320
211 gccggcgatc cggcgctggc ctacggctac gcccaggacc aggaaccgca cgcacgcggc      1380
213 cggatccgca acggtgccct gctgcgggtc tatgtgccgc gctcgagcct gccgggcttc      1440
215 taccgcacca gcctgaccct ggccgcgccg gaggcggcgg gcgaggtcga acggtgatc      1500
217 ggccatccgc tgccgtcgcg cctggacgcc atcaccggcc ccgaggagga aggcgggcgc      1560
219 ctggagacca ttctcggtcg gccgctggcc gagcgaccgc tgggtgattcc ctcggcgatc      1620
221 cccaccgacc cgcgcaacgt cggcggcgac ctcgaccgtt ccagcatccc cgacaaggaa      1680
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225 ctgaagtaa
228 <210> SEQ ID NO: 4
229 <211> LENGTH: 672
230 <212> TYPE: DNA
231 <213> ORGANISM: pMC75 plasmid coding sequence
233 <400> SEQUENCE: 4
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236 tccatctctt gcagatctag tcagatcatt gtacatagta atggaaacac ctatttagaa      120
238 tggtagctgc agaaaccagg ccagttctcca aagctcctga tctacaaagt ttccaaccga      180
240 ttttctgggg tcccagacag gttcagtggc agtggatcag ggacagattt cacactcaag      240
242 atcagcagag tggaggctga ggatctggga gtttattact gctttcaagg ttcacatgtt      300
244 ccattcacgt tcggctcggg gacaaagttg gaaataaaac gggctgatgc tgcaccaact      360
246 gtatccatct tcccaccatc cagtgagcag ttaacatctg gaggtgcctc agtcgtgtgc      420
248 ttcttgaaca acttctaccc caaagacatc aatgtcaagt ggaagattga tggcagtga      480
250 cgacaaaatg gcgtcctgaa cagttggact gatcaggaca gcaaagacag cacctacagc      540
252 atgagcagca cctcaccgtt gaccaaggac gagtatgaac gacataacag ctatacctgt      600
254 gaggccactc acaagacatc aacttcaccc attgtcaaga gcttcaacag gaatgagtgt      660
256 ggtaaagctt aa

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TIME: 09:00:14

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Output Set: N:\CRF4\01102006\J562627.raw

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289 gaactcctgg ggggaccgtc agtcttctc tccccccaa aacccaagga caccctcatg      780
291 atctcccggg cccctgaggt cacatgcgtg gtggtggacg tgagccacga agaccctgag      840
293 gtcaagttca actggtacgt ggacggcgtg gaggtgcata atgccaaag aaagccgcgg      900
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297 tggctgaatg gcaaggagta caagtgcaag gtctccaaca aagccctccc agcccccac      1020
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311 ggtggtggcg gttctaaagc ttccggaggt cccgagggcg gcagcctggc cgcgctgacc      1440
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325 ggcgcgaggt tcctcggcga cggcggcgac gtcagcttca gcacccgcgg cacgcagaac      1860
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337 accctggccg cgccggaggc ggcgggcgag gtcgaacggc tgatcggcc tccgtgccg      2220
339 ctgcgcctgg acgccatcac cgccccgag gaggaaggcg ggcgctgga gaccattctc      2280
341 ggctggccgc tggccgagcg caccgtggtg attccctcgg cgatccccac cgaccgcgc      2340
343 aacgtcggcg gcgacctcga cccgtccagc atccccgaca aggaacaggc gatcagcgcc      2400
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350 <212> TYPE: DNA

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## RAW SEQUENCE LISTING

DATE: 01/10/2006

PATENT APPLICATION: US/10/562,627

TIME: 09:00:14

Input Set : N:\DA\PTO.DA.txt

Output Set: N:\CRF4\01102006\J562627.raw

351 <213> ORGANISM: pKL4 plasmid coding sequence  
353 <400> SEQUENCE: 6

354	atgcatcacc	atcaccatca	cgatgtgaag	ctggtggaat	ctggaggagg	cttagtgcag	60
356	cctggagggt	ccctgaaact	ctcctgtgca	acctctggat	tcactttcag	tgactattac	120
358	atgtattggg	ttcgccagac	tccagagaag	aggctggagt	gggtcgcata	cattagtaat	180
360	gatgatagtt	ccgccgctta	ttcagacact	gtaaagggcc	gggttcaccat	ctccagagac	240
362	aatgccagga	acaccctcta	cctgcaaatg	agccgtctga	agtctgagga	cacagccata	300
364	tattcctgtg	caagaggact	ggcctgggga	gcctggtttg	cttactgggg	ccaagggact	360
366	ctggtcactg	tctctgcagc	caaaacgaca	ccccatctg	tctatccact	ggccccctga	420
368	tctgctgccc	aaactaactc	catggtgacc	ctgggatgcc	tggtcaaggg	ctattttccct	480
370	gagccagtga	cagtgcctg	gaactctgga	tccctgtcca	gcggtgtgca	caccttccca	540
372	gctgtcctgc	agtctgacct	ctacactctg	agcagctcag	tgactgtccc	ctccagcacc	600
374	tggcccagcg	agaccgtcac	ctgcaacgtt	gcccacccgg	ccagcagcac	caaggtggac	660
376	aagaaaattg	tgcccagggg	ttgtggtgct	aagccttgca	tagctacaca	agcttccggg	720
378	ggtggcggat	ctggagggtg	cggaagcgga	ggtcccagg	tgacaggggg	aatggcaagc	780
380	aagtgggatc	agaaggggat	ggacattgcc	tatgaggagg	cggccttagg	ttacaaagag	840
382	ggtggtgttc	ctattggcgg	atgtcttatc	aataacaaag	acggaagtgt	tctcggtcgt	900
384	ggtcacaaca	tgagatttca	aaagggatcc	gccacactac	atggtgagat	ctccactttg	960
386	gaaaactgtg	ggagattaga	gggcaaagt	tacaaagata	ccactttgta	tacgacgctg	1020
388	tctccatgcg	acatgtgtac	aggtgccatc	atcatgtatg	gtattccacg	ctgtgttgct	1080
390	ggtgagaacg	ttaatttcaa	aagtaagggc	gagaaatatt	tacaaactag	aggtcacgag	1140
392	gttgtgtgtg	ttgacgatga	gaggtgtaaa	aagatcatga	aacaatttat	cgatgaaaga	1200
394	cctcaggatt	ggtttgaaga	tattggtgag	tag			1233

397 <210> SEQ ID NO: 7  
398 <211> LENGTH: 4871  
399 <212> TYPE: DNA

400 <213> ORGANISM: pMC74 plasmid full sequence  
402 <400> SEQUENCE: 7

403	taatacgact	cactataggg	agaccacaac	ggtttcctc	tagaaataat	tttgtttaac	60
405	tttaagaagg	agatatacat	atggatgtga	agctggtgga	atctggagga	ggcttagtgc	120
407	agcctggagg	gtccctgaaa	ctctcctgtg	caacctctgg	attcactttc	agtgactatt	180
409	acatgtattg	ggttcgccag	actccagaga	agaggctgga	gtgggtcgca	tacattagta	240
411	atgatgatag	ttccgccgct	tattcagaca	ctgtaaaagg	ccggttcacc	atctccagag	300
413	acaatgccag	gaacaccctc	tacctgcaaa	tgagccgtct	gaagtctgag	gacacagcca	360
415	tatattcctg	tgcaagagga	ctggcctggg	gagcctggtt	tgcttactgg	ggccaagggg	420
417	ctctggtcac	tgtctctgca	gccccaaacg	cacccccatc	tgtctatcca	ctggccccctg	480
419	gatctgctgc	ccaaactaac	tccatggtga	ccctgggatg	cctggtcaag	ggctatttcc	540
421	ctgagccagt	gacagtgacc	tggaactctg	gatccctgtc	cagcgtgtg	cacaccttcc	600
423	cagctgtcct	gcagtctgac	ctctacactc	tgagcagctc	agtgactgtc	ccctccagca	660
425	cctggcccag	cgagaccgtc	acctgcaacg	ttgcccaccc	ggccagcagc	accaaggtgg	720
427	acaagaaaat	tgtgcccagg	gattgtggta	gtaagcctag	cataagtaca	aaagcttccg	780
429	gaggtcccga	gggcggcagc	ctggccgcgc	tgaccgcgca	ccaggcttgc	cacctgccgc	840
431	tggagacttt	caccgcgtcat	cgccagccgc	gcggtcggga	acaactggag	cagtgcggct	900
433	atccggtgca	gcggtcggtc	gccctctacc	tggcggcgcg	gctgtcgtgg	aaccaggtcg	960
435	accaggtgat	ccgcaacgcc	ctggccagcc	ccggcagcgg	cggcgacctg	ggcgaagcga	1020
437	tccgcgagca	gccggagcag	gcccgtctgg	ccctgaccct	ggccgcggcc	gagagcgagc	1080
439	gcttcgtccg	gcagggcacc	ggcaacgacg	aggccggcgc	ggccaacggc	ccggcggaca	1140
441	gcggcgacgc	cctgctggag	cgcaactatc	ccactggcgc	ggagtctctc	ggcgacggcg	1200
443	gcgacgtcag	cttcagcacc	cgcggcacgc	agaactggac	ggtggagcgg	ctgctccagg	1260

FYI

The type of errors shown exist throughout  
the Sequence Listing. Please check subsequent  
sequences for similar errors.

**VERIFICATION SUMMARY**

DATE: 01/10/2006

PATENT APPLICATION: US/10/562,627

TIME: 09:00:15

Input Set : N:\DA\PTO.DA.txt

Output Set: N:\CRF4\01102006\J562627.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date